

# United States Patent and Trademark Office

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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO 4300				
10/642,946	08/18/2003	James W. Ryan	JR-14000-CON					
75	90 10/17/200	i	EXAMINER					
Cheryl H. Agr P.O. Box 806	is, Ph.D.		ZARA,	JANE J				
	0803		ART UNIT	PAPER NUMBER				
			1635					
		DATE MAILED: 10/17/2006						

Please find below and/or attached an Office communication concerning this application or proceeding.

	Application No.	Applicant(s)							
	10/642,946	RYAN, JAMES W.							
Office Action Summary	Examiner	Art Unit							
	Jane Zara	1635							
The MAILING DATE of this communication app Period for Reply	ears on the cover sheet with the c	orrespondence address							
A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA  - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication.  - If NO period for reply is specified above, the maximum statutory period w  - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be time will apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE!	N. nely filed the mailing date of this communication. D (35 U.S.C. § 133).							
Status									
<ol> <li>Responsive to communication(s) filed on <u>07 September 2006</u>.</li> <li>This action is <b>FINAL</b>. 2b) This action is non-final.</li> <li>Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under <i>Ex parte Quayle</i>, 1935 C.D. 11, 453 O.G. 213.</li> </ol>									
Disposition of Claims									
4) Claim(s) 1-4,6-16 and 23-26 is/are pending in t 4a) Of the above claim(s) 7,9,12,13 and 23-26 i  5) Claim(s) is/are allowed.  6) Claim(s) 1-4,6,8,10,11 and 14-16 is/are rejecte  7) Claim(s) is/are objected to.  8) Claim(s) are subject to restriction and/or  Application Papers	s/are withdrawn from considerati  d.  election requirement.	on.							
9) The specification is objected to by the Examiner.  10) The drawing(s) filed on is/are: a) accepted or b) objected to by the Examiner.  Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).  Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).  11) The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.  Priority under 35 U.S.C. § 119  12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).  a) All b) Some * c) None of:  1. Certified copies of the priority documents have been received.  2. Certified copies of the priority documents have been received in Application No  3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).									
* See the attached detailed Office action for a list of the section fo	of the certified copies not receive	d.							
Attachment(s)  1) Notice of References Cited (PTO-892)  2) Notice of Draftsperson's Patent Drawing Review (PTO-948)  3) Information Disclosure Statement(s) (PTO/SB/08)  Paper No(s)/Mail Date 1-29-04.	4) Interview Summary Paper No(s)/Mail Da 5) Notice of Informal Pa	nte							

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## **DETAILED ACTION**

This Office action is in response to the communication filed 9-7-06.

Claims 1-4, 6-16, 23-26 are pending in the instant application.

#### Election/Restrictions

Applicant's election with traverse of Group I, claims 1-4, 6, 8, 10, 11 and 14-16, and SEQ ID No. 6 in the reply filed on 9-7-06 is acknowledged. The traversal is on the ground(s) that Groups II-IV are ultimately linked to the claims in Group I and are subject to the right of rejoinder one there is indication of allowable subject matter of Group I. Applicant also argues that no undue burden would exist to search all of the sequences originally claimed in addition to the elected sequence. This is not found fully persuasive because the searches required for proper examination of all of the sequences and all of the different methods claimed would be burdensome to the examiner and, even though the searches of the prior art and appropriate data bases for some of the sequences and groups might be overlapping, they would not be coextensive. Applicant is correct that the linking claims would be rejoined upon allowance of the elected Group, provided that the methods were commensurate in scope with the allowed composition claims.

The requirement is still deemed proper and is therefore made FINAL.

Claims 7, 9, 12, 13, 23-26 are withdrawn from further consideration pursuant to 37 CFR 1.142(b), as being drawn to a nonelected invention, there

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being no allowable generic or linking claim. Applicant timely traversed the restriction (election) requirement in the reply filed on 9-7-06.

## Claim Rejections - 35 USC § 112

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

Claims 1-4, 6, 8, 10, 11 and 14-16 are rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the written description requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

The claims are drawn to compositions comprising isolated genomic nucleic acid molecules having a nucleotide sequence at least 95% identical to any polynucleotide encoding the polypeptide of SEQ ID NO. 3, or any nucleic acid molecule that hybridizes to these polynucleotides, and isolated nucleic acids comprising at least 20 nucleotides that hybridize with high stringency to an intronic region of SEQ ID No. 6.

The specification, claims and the art do not adequately describe the distinguishing features or attributes concisely shared by the members of the genus comprising these nucleic acids that are at least 95% homologous to any polynucleotide that encodes the polypeptide of SEQ ID NO. 3, or nucleic acids

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that specifically hybridize to them, or any polynucleotide that stringently hybridizes to the intronic sequences of SEQ ID No. 6. The specification teaches the sequence of SEQ ID No. 6, encoding the polypeptide of SEQ ID NO. 3. The genus of nucleic acids claimed, however, encompasses a myriad of structures (e.g. thousands and thousands of nucleic acid sequences) and the specification and claims do not adequately teach a representative number of species for the broad genus claimed, and which provide for the function claimed, of having human adipocyte enhancer binding protein 1 activity.

Concise structural features that could distinguish structures within the genus from others are missing from the disclosure. No common structural attributes identify the members of the claimed genus, and distinguish members within the claimed genus from those outside of it, and which provide for the function claimed, of having human adipocyte enhancer binding protein 1 activity. One of skill in the art would reasonably conclude that the disclosure fails to provide a representative number of species to describe the genus claimed.

### Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

<sup>(</sup>b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

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(e) the invention was described in (1) an application for patent, published under section 122(b), by another filed in the United States before the invention by the applicant for patent or (2) a patent granted on an application for patent by another filed in the United States before the invention by the applicant for patent, except that an international application filed under the treaty defined in section 351(a) shall have the effects for purposes of this subsection of an application filed in the United States only if the international application designated the United States and was published under Article 21(2) of such treaty in the English language.

Claims 1-4, 6 and 10 are rejected under 35 U.S.C. 102(b) as being anticipated by Layne et al.

Layne et al (J. Biol. Chem., Vol. 273, No. 25, pages 15,654-15,660, 1998) teach compositions and methods of recombinant polypeptide expression in vitro comprising the nucleic acid encoding the polypeptide of SEQ ID NO. 3 and further comprising a carrier (e.g. water), as well as an expression vector and recombinant host cells comprising the nucleic acid encoding the polypeptide of SEQ ID No. 3 (see the abstract on p. 15,654, text on p. 15,655, fig. 1 on p. 15,656, discussion on pp. 15,658-9. See also the accompanying alignment between SEQ ID No. 3 of the instant application and AF053944 of Layne et al).

Claims 8 and 11 are rejected under 35 U.S.C. 102(b) as being anticipated by Noonberg et al.

Noonberg et al (USPN 5,624,803) teach a compositions comprising a polynucleotide at least 20 nucleobases in length that specifically hybridizes under stringent condition with an intronic sequence of SEQ ID NO. 6 and further comprising a carrier, water (see the accompanying sequence alignment data between nucleotides 3002-3237 of SEQ ID NO. 6 and SEQ ID No. 20 of

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Noonberg et al. In the alignment data, nucleotide no. 3002 of SEQ ID No. 6 = nucleotide no. 1).

Claims 8 and 11 are rejected under 35 U.S.C. 102(e) as being anticipated by Denney et al.

Denney et al (USPN 5,972,334) teach a compositions comprising a polynucleotide at least 20 nucleobases in length that specifically hybridizes under stringent condition with an intronic sequence of SEQ ID NO. 6 and further comprising a carrier, water (see the accompanying sequence alignment data between nucleotides 1967-2208 of SEQ ID NO. 6 and SEQ ID No. 35 of Denney et al. In the alignment data, Nucleotide no. 3002 of SEQ ID No. 6 = nucleotide no. 1).

#### Conclusion

Certain papers related to this application may be submitted to Art Unit 1635 by facsimile transmission. The faxing of such papers must conform with the notices published in the Official Gazette, 1156 OG 61 (November 16, 1993) and 1157 OG 94 (December 28, 1993) (see 37 C.F.R. ' 1.6(d)). The official fax telephone number for the Group is **571-273-8300**. NOTE: If Applicant *does* submit a paper by fax, the original signed copy should be retained by applicant or applicant's representative. NO DUPLICATE COPIES SHOULD BE SUBMITTED so as to avoid the processing of duplicate papers in the Office.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to **Jane Zara** whose telephone number is **(571)** 

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272-0765. If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Peter Paras, can be reached on (571) 272-4517. Any inquiry regarding this application should be directed to the patent analyst, Katrina Turner, whose telephone number is (571) 272-0564. Any inquiry of a general nature or relating to the status of this application should be directed to the Group receptionist whose telephone number is (703) 308-0196.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

Jane Zara 10-10-06

JANE ZARA, PH.D.

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I.T., Perrella, M.A., Blanar, M.A., Haber, E. and Lee, M.E.
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M.D., Endege,W.O., Jain,M.K., Yet,S.-F., Hsieh,C.-M.-E.
Submission
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: Gla-MRR-1989 Cardiovascular Biology Laboratory, Harvard
ited (16-MRR-1989) Cardiovascular Biology Laboratory, Harvard
of Public Health, 677 Huntington Ave, Boston, MA 02115, USA
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Ov 421 ThrGlyAlaThrG	1400	27 441 InrGinTrplied 	2y 461 GlyArgAspSerS 	2y .481 AspSerGlnThrT 	11	Oy 521 ArgileTyrProf. 	541	Oy 561 AspPheArgHisH 	581		2000	2060	2120	2180	2240	2300	2360	0y 761 AsnLeuAsnGlyG 
260 TTCCTGTCAGAGCTAGAACCTGAGCCCCGGGAGGACGACGTGGAGGCCCCCGCCGCCTCCC 319 Ov	61 GluProThrProArgValArgLy8AlaGlnAlaGlyGlyLy8ProGlyLy8ArgProGly 80	81 ThralaalagluValProProGluLysThrLysAspLysGlyLysLysEysGlyLysLysBsp 100	LysGlyProLysValProLysGluSerLeuGluGlySerProArgProLysLysGly	LysGluLysProProLysAlaThrLysLysProLysGluLysProProLysAlaThrLys		Oy  161 ThrLysLysProProSerGlyLysArgProProIleLeuAlaProSerGluThrLeuGlu 180	TrpProLeuProProProProGerProGluGluLeuProGlnGluGlyGlyAla 200	ProLeuSerAsnAsnTrpGlnAsnProGlyGluGluThrHisValGluAlaGlnGluHis 220	GinProGluProGluGluThrGluGlnProThrLeuAspTyrAsnAspGlnIleGlu 240	241 ArgGluAepTyxGluAepPheGluTyrIleArgArgGlnLysGlnProArgProProP 260	280 979	281 ProAlaProGludluArgIleGluProProValLysProLeuLeuProProLeuProPro 300	OY 301 ASPTYTG1YASPG1YTYTVA111eProAsnTYTASPASPMetASPTYTTYTPTPHG1YPro 320 1040 GACTATGGTGATGGTTACGTGATCCCCAACTACGATGACATGACTATTACTTTGGGCCT 1099	OY  321 ProProProGlnLysProAspAlaGluArgGlnThrAspGluGluLysGluGluLeuLys 340	OY  341 LysProLysGluAspSerSerProLysGluGluThrAspLysTrpAlaValGluLys 360	Oy  361 GlyLysAapHisLysGluProArgLysGlyGluGluLeuGluGluGluGluTrpThrProThr 380  1220 GGCAAGGACCACAAAAGAGCCCCGAAAGGGCGAGGAGTGGAGGAGGAGGAGGAGGACCTACG 1279	381 GluLy8ValLy8CySProProIleGlyMetGluSerHisArgIleGluAspAsnGlnIle 400	401 ArgAlaSerSerMetLeuArgHisGlyLeuGlyAlaGlnArgGlyArgLeuAsnMetGln 420   Dy   ArgAlaSerSerMetLeuArgHisGlyLeuGlyAlaGlnArgGlyArgLeuAsnMetGln 420   Did   D
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ThrProvalLeuSerGluLeuProGluProvalvalAlaArgPheIle 520 LeuThrTrpAsnGlySerLeuCysMetArgLeuGluValLeuGlyCys 540 ValtyrSertyrTyrAlaGlnAsnGluValValAlaThrAspAspLeu 560 AsnàsnleuprolleprogluargTyrleuSerproaspalaThrVal GlubspasptyrtyraspGlyalaTrpCysAlaGlubspAspAlaArg 

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NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
Email: cgapbs-remail.nih.gov
Tissue Procurement: Life Technologies, Inc.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: National Institutes of Health Intramural
Sequencing Center (NISC),
Gaithersburg, Maryland;
Web site: http://www.nisc.nih.gov/
Contact: nisc.mgc@nhgri.nih.gov/
Contact: nisc.mgc@nhgri.nih.gov/
Contact: N., Ayele,K., Beckstrom-Sternberg,S.M., Benjamin,B.,
Blakesley,R.W., Bouffard,G.G., Breen,K., Brinkley,C., Brooks,S.,
Dietrich,N.L., Granite,S., Guan,X., Gupta,J., Haghighi,P.,
Hansen,N., Ho,S.-L., Karlins,E., Kwong,P., Laric,P., Legaspi,R.,
Mcduro,Q.L., Masiello,C., Maskeri,B., Mastrian,S.D., McCloskey,J.C.,
McDowell,J., Pearson,R., Stantripop,S., Thomas,P.J., Touchman,J.W.,
TBURGGON,C., Vogt,J.L., Walker,M.A., Wetherby,K.D., Wiggins,L.,
Young,A., Zhang,L.-H. and Green,E.D.
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Strausberg, L. Peingold, B.A., Grouse, L.H., Derge, J.G., Klausner, R.D., Colling, P.S., Wagner, L., Shenmen, C.M., Schuler, G. Klausner, R.D., Colling, P.S., Wagner, L., Shenmen, C.R., Schuler, G.D., Altechul, S.P., Zeeberg, B., Buetow, K.H., Schaefer, C.F., Bhat, N.K., Hopkins, R.F., Jordan, H., Moore, T., Max, S.I., Wang, J., Hsieh, F., Diatchenko, L., Marusina, K., Parmer, A.A., Rubin, G.M., Hong, L., Stapleton, M., Soares, M.B., Bonaldo, M.F., Casavant, T.L., Schetz, T.E., Brownstein, M.J., Usdin, F.B., Toshiyuki, S., Schetz, T.E., Brownstein, M.J., Usdin, F.B., Toshiyuki, S., Abramson, R.D., Mullahy, S.J., Bosak, S.A., McEwan, P.J., Mare, G.J., Malek, J.A., Gunaratne, P.H., Richards, S., Worley, K.C., Hale, S., Garcia, A.M., Gay, L.J., Hulyk, S.M., Worley, K.C., Hale, S., Garcia, A.M., Madan, A., Rodrigues, S., Sanchez, A., Whiling, M., Madan, A., Rodrigues, S., Sanchez, A., Whiling, M., Madan, A., Young, A.C., Shevchenko, Y. Butfeard, G.G., Blakesley, R.W., Puckeon, M.C., Rodriguez, A.C., Grimwood, J., Schmutz, J., Myers, R.M., Butterfield, Y.S., Krzywinski, M.I., Skalska, U., Smallus, D.E., Schnerch, A., Schein, J. B., And Marra, M.A.
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                                                                                                                             linear PRI 30-JUN-2004
(cDNA clone MGC:46180
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This clone was selected for full length sequencing because it passed the following selection criteria: matched mRNA gi: 4755145.
Location/Qualifiers
1. .4125
                                                                                                                                                                                                                                                                                                                                  Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
Submitted (15-007-2002) National Institutes of Health, Mammalian
Gene Collection (MGC), Cancer Genomics Office, National Cancer
Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
3560 CAGGCATTCCCCTTCACAACAGTAGAGACCTACACAGTGAACTTTGGGGACTTC 3613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Proc. Natl. Acad. Sci. U.S.A. 99 (26), 16899-16903 (2002)
                                                                                                                                mRNA
mRNA
                                                                                                                           4125 bp
Homo sapiens AE binding protein 1,
IMAGE:5764769), complete cds.
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KEYWORDS
SOURCE
ORGANISM
                                                                                                                                                        DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE
AUTHORS
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REFERENCE
AUTHORS
TITLE
                                                                                                                                                                                                         ACCESSION
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                                                                                 RESULT 4
BC038588
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APPLICANT: Woodward, Robert
APPLICANT: Woodward, Robert
APPLICANT: Woodward, Robert
APPLICANT: Ly, NGCC
APPLICANT: Ly, NGCC
APPLICANT: Ly, NGCC
TITLE OF INVENTION: CHENDIC AND COMPOSITIONS FOR DIAGNOSING AND MONITORING AUTOIMMUN
TITLE OF INVENTION: CHENDIC 100
FILE REFERENCE: 506612000120
CURRENT APPLICATION NUMBER: US 10/006, 290
PRIOR PILING DATE: 2001-10-22
PRIOR FILING DATE: 2001-10-22
PRIOR FILING DATE: 2001-10-22
PRIOR FILING DATE: 2001-10-22
PRIOR FILING DATE: 2001-10-33
PRIOR FILING DATE: 2001-10-32
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0
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8.8%; Score 20.8; DB 2;
Best Local Similarity 78.1%; Pred. No. 3.2e+03;
Matches 25; Conservative 0; Mismatches 7;
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Patent No. 6905827
                                                                                                                                                                                                                                                                        Oy 3, 2 ... 137 CACCTGCTCCCCTTCTCCT
  TYPE: nucleic acid

TYPE: nucleic acid

STRANDEDNESS: double

TOPOLOGY: linear
US-08-324-001-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Wohlgemuth, Jay
APPLICANT: Fry, Kirk
APPLICANT: Woodward, Robert
SEQUENCE CHARACTERISTICS LENGTH: 43 base pairs
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US-10-131-827-584
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Pred. No. 3.26+03;
0; Mismatches 7; Indels
                                                                                                                               COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/324,001
FILING DATE: 13-0CT-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MONROY, GLADYS H.
REGISTRATION NUMBER: 32,430
REFERENCE/DOCKET NUMBER: 22000-20544.20
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 813-5600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: 706141 MRSN FORESSFO INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8.8%;
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Best Local Similarity 78.1
Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 43 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                           ZIP: 94304-1018
COMPUTER READABLE FORM:
                                     USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
US-08-324-001-19
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APPLICANT: Wohlgemuth, Jay
APPLICANT: Fry, Kirk
APPLICANT: Pry, Kirk
APPLICANT: Woodward, Robert
APPLICANT: Woodward, Robert
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING
TITLE OF INVENTION: TRANSPLANT REJECTION
FILE REFERENCE: 506610121
CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT PILING DATE: 2002-08-05
                                                      Gaps
                                                         ö
                                                                                                       71 GTGAGAGCTGGTTTGAAAGAGCCTGGCCCCCTCTCATTC 112
                                                                                                                                                      Gregacccrerrierrraaararrererrecearereare 47
  Score 19.6; DB 3;
Pred. No. 8.1e+03;
0; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 9190
SOFTWARE: Patentin version 3.1
SEQ ID NO 584
LENGTH: 50
                                                                                                                                                                                                                                                                   584, Application US/10131831
o. 7026121
Query Match
Best Local Similarity 66.79
Marches 28; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IMP PC compatible
COMPUTER: IMP PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/324,001
FILING DATE: 13-OCT-1994
CLASSIFICATION: 435

FILING DOLL 435
CIASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MONROY, GLADYS H.

ELECOMMUNICATION INFORMATION

REFERENCE/DOCKET NUMBER:

(415) 813-5600

INFORMATION FOR SEQ ID NO:

Length 50;

TYPE: DNA
CRGANISM: Homo sapiens
US-10-131-827-584

IN VIVO OLIGONUCLEOTIDE GENERATOR, AND METHODS OF TESTING THE BINDING AFFINITY OF TRIPLEX FORMING OLIGONUCLEOTIDES DERIVED THEREFROM

Sequence 20, Application US/08324001 Patent No. 5624803

GENERAL INFORMATION:
APPLICANT: NOONBERG, SARAH B.
APPLICANT: HUNT, C. ANTHONY

APPLICANT: HUNT, C. ANT TITLE OF INVENTION: IN TITLE OF INVENTION: MET TITLE OF INVENTION: OLI NUMBER OF SEQUENCES: 25 CORRESPONDENCE ADDRESS:

E: MORRISON & FOERSTER 755 PAGE MILL ROAD

STREET: 755 TTY: PALO ALTO

ADDRESSEE:

USA

COUNTRY:

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                                                                                                                                                                                                                                                                                                            APPLICANT: LY, NGOC
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING
TITLE OF INVENTION: TRANSPLANT REJECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               50 crerececrirececririca di riva de la consecrire de la c
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COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/644,664B
FILING DATE: 01-MAY-1996
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: Ingolia, Diane R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               35 CTGGGAGCTCCACCTCCCCAGCTGCTGCCTCACCTGCTTTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 19.6; DB 5;
Pred. No. 4.5e+03;
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GENERAL INFORMATION:
APPLICANT: Denney Jr., Dan W.
TITLE OF INVENTION: Gene Amplication Methods
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Medlen & Carroll, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      : 220 Montgomery Street, Suite 2200
San Francisco
California
                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 506612000121
CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT FILING DATE: 2002-08-05
PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
                                                        Application US/10131831
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REFERENCE/DOCKET NUMBER: GENI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 9190
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                     Fry, Kirk
Woodward, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 397-8338 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                         GENERAL INFORMATION:
APPLICANT: Wohlgemuth, Jay
APPLICANT: Fry, Kirk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 28; Conserv
US-10-131-831-1159/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 7
US-08-644-664B-35/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 1159
LENGTH: 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE:
                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Length 39;

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Robert L.

CLONING AND DEVELOPMENTAL EXPRESSION OF

PEA RIBULOSE-1,5-BISPHOSPHATE CARBOXYLASE/OXYGENASE LARGE

BEA RIBULASE-1,5-BISPHOSPERASE

SUBUNIT N-METHYLIRANSFERASE
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                                                   Gaps
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                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Denney Jr., Dan W.
TITLE OF INVENTION: Vaccines For Treatment Of Lymphoma And
TITLE OF INVENTION: Leukemia
NUMBER OF SEQUENCES: 80
CORRESPONDENCE ADDRESS:
ADDRESSEE: Medien & Carroll, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 39;
                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION OF E. US/08/761,277A
FILING DATE: 06-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 19.2; DB 2;
Pred. No. 5.6e+03;
0; Mismatches 8;
  7.9%; Score 19.2; DB 2; 75.0%; Pred. No. 5.6e+03; tive 0; Mismatches 8;
                                                                                         211 TGTATCAGATACCCAACTAAAACTGGATTCAC 242
                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Medlen & Carroll, LLP
STREET: 220 Montgomery Street, Suite 2200
CITY: San Francisco
STATE: California
COUNTRY: United States Of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     211 TGTATCAGATACCCAACTAAAACTGGATTCAC 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  readacadraceadecradacerregarecae 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: MacKright, Kamrin T.
REGISTRATION NUMBER: 38,230
REFERENCE/DOCKET NUMBER: GENITOPE-02406
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
                                                                                                                                     37 TGAAACAGATACGAACCTAAACTTGGATCCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 424
RRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/644,664
FILING DATE: 01-MAY-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                             Sequence 35, Application US/08761277A Patent No. 5972334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25, Application US/08391000
Patent No. 5723752
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 75.0%;
Matches 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 397-8338
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 39 base pairs
TYPE: nucleic acid
                                              24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: HOUTZ, RO
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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Query Match
Best Local Similarity
Matches 24, Conserva
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MOLECULE TYPE:
                                                                                                                                                                                                     RESULT 8
US-08-761-277A-35/C
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MOLECULE TYPE: DNA (genomic)

US-08-644-664B-35

linear